# **AMENDMENTS TO THE CLAIMS**

1. (currently amended): An isolated chimeric protein having the enzymatic activity of a nucleotidase, which chimeric protein comprises, from N-terminus to C-terminus:

- a) a first peptidyl fragment comprising a bacterial leader sequence comprising an amino acid sequence having at least 80% identity with the amino acid sequence as set forth in SEQ ID NO:1;
- b) a second peptidyl fragment <u>comprising</u> that binds to an antibody that specifically binds to an <u>the</u> amino acid sequence as set forth in SEQ ID NO:2 <u>or a derivative thereof having a conservative amino acid substitution where the derivative retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and</u>
- c) a third peptidyl fragment comprising an amino acid sequence having at least 80% identity with the amino acid sequence as set forth in SEQ ID NO:3.

# 2-11. (canceled)

12. (previously presented): The isolated chimeric protein of claim 1, wherein the first and second peptidyl fragments are linked via a cleavable linkage.

# 13-20. (canceled)

- 21. (previously presented): The isolated chimeric protein of claim 1, which further comprises, at its C-terminus a fourth peptidyl fragment comprising a peptide tag.
- 22. (previously presented): The isolated chimeric protein of claim 21, wherein the peptide tag is selected from the group consisting of FLAG, HA HA1, c-Myc, 6-His, AU1, EE, T7, 4A6,  $\epsilon$ , B, gE, and Ty1 tag.

23. (currently amended): The An isolated chimeric protein of claim 1, which comprises the amino acid sequence set forth in SEQ ID NO:4 (mggsgddddlalALERELLVATQAVRKASLLTKRIQSEVISHKDSTTITKNDNSPVTTGDYAAQT IIINAIKSNFPDDKVVGEESSSGLSDAFVSGILNEIKANDEVYNKNYKKDDFLFTNDQFPLKS LEDVRQIIDFGNYEGGRKGRFWCLDPIDGTKGFLRGEQFAVCLALIVDGVVQLGCIGCPNL VLSSYGAQDLKGHESFGYIFRAVRGLGAFYSPSSDAESWTKIHVRHLKDTKDMITLEGVEK GHSSHDEQTAIKNKLNISKSLHLDSQAKYCLLALGLADVYLRLPIKLSYQEKIWDHAAGNV IVHEAGGIHTDAMEDVPLDFGNGRTLATKGVIASSGPRELHDLVVSTSCDVIQSRNAkgelegl pipnpllrtghhhhhh).

# 24-30. (canceled)

- 31. (currently amended): A method for assaying for sodium ions in a sample, which method comprises:
- a) contacting the sample with <u>a</u> the chimeric protein-of claim 1, comprising, from N-terminus to C-terminus:
  - (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
  - (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution where the derivative retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
  - (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a sodium-sensitive 3'(2'),5'-bisphosphate nucleotidase, wherein the nucleotidase consumes adenosine 3',5'-bisphosphate (PAP) and forms AMP and P<sub>i</sub>; and

b) assessing the consumption of PAP or the formation of AMP or  $P_i$  in step a) to determine the presence or amount of sodium ions in the sample.

- 32. (original): The method of claim 31, wherein the sample is a biological sample.
- 33. (original): The method of claim 32, wherein the biological sample is a blood sample.
- 34. (original): The method of claim 33, wherein the blood sample is a plasma, serum, red blood cell, or whole blood sample.

# 35-36. (canceled)

- 37. (original): The method of claim 31, wherein the amount of AMP formed is inversely related to the amount of sodium ions in the sample.
- 38. (original): The method of claim 31, which is used in prognosis or diagnosis of a disease or disorder.
- 39. (currently amended): A method for assaying for sodium ions in a sample, which method comprises:
- a) contacting the sample with a first composition comprising adenosine 3',5'-bisphosphate (PAP);

b) contacting the sample with a second composition comprising <u>a</u> the-chimeric protein of claim 1, comprising, from N-terminus to C-terminus:

- (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
- (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution that retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
- (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a sodium-sensitive 3'(2'),5'-bisphosphate nucleotidase; and

- c) assessing the production of AMP to determine the presence or amount of sodium ions in the sample.
  - 40. (original): The method of claim 39, wherein the sample is a biological sample.
  - 41. (original): The method of claim 40, wherein the biological sample is a blood sample.
- 42. (original): The method of claim 41, wherein the blood sample is a plasma, serum, red blood cell, or whole blood sample.
  - 43. (canceled)
- 44. (original): The method of claim 39, wherein the first composition further comprises 4-aminoantipyrine (4-AA), N-ethyl-N-(2-hydroxy-3-sulfopropyl)-3-m-toluidine (EHSPT), purine nucleoside phosphorylase, xanthine oxidase, and peroxidase, and the second composition further comprises adenosine deaminase, 5'-nucleotidase, and MgCl<sub>2</sub>.

45. (currently amended): A kit for assaying for sodium ions in a sample, which kit comprises

- a) a first composition comprising <u>a</u> the chimeric protein <del>of claim 1, comprising, from N-terminus to C-terminus:</del>
  - (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
  - (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution that retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
  - (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a sodium-sensitive 3'(2'),5'-bisphosphate nucleotidase that consumes adenosine 3',5'-bisphosphate and forms AMP and  $P_i$ ; and

- b) means for assessing the product formed or the substrate consumed by the nucleotidase to determine the presence or amount of the sodium ions in the sample.
- 46. (original): The kit of claim 45, wherein the first composition further comprises adenosine deaminase, 5'-nucleotidase and MgCl<sub>2</sub>.
- 47. (previously presented): The kit of claim 45, further comprising a second composition comprising 4-aminoantipyrine (4-AA), N-ethyl-N-(2-hydroxy-3-sulfopropyl)-3-m-toluidine (EHSPT), purine nucleoside phosphorylase, xanthine oxidase, and peroxidase, wherein the reaction of 4-AA and EHSPT in the presence of peroxidase is the means for assessing the product formed.
- 48. (original): The kit of claim 45, which further comprises a low sodium serum standard and a high sodium serum standard.

6

sd-448989

- 49. (canceled)
- 50. (currently amended): A method for assaying for lithium ions in a sample, which method comprises:
- a) contacting the sample with <u>a</u> the chimeric protein <del>of claim 1, comprising, from N-terminus to C-terminus.</del>
  - (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
  - (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution that retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
  - (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a lithium-sensitive 3'(2'),5'-bisphosphate nucleotidase, wherein the nucleotidase consumes adenosine 3',5'-bisphosphate (PAP) and forms AMP and P<sub>i</sub>; and

- b) assessing the amount of PAP consumed or AMP or Pi formed in step (a) to determine the presence or absence of lithium ions in the sample.
- 51. (original): The method of claim 50 further comprising first contacting the sample with a sodium blocking agent.
- 52. (original): The method of claim 51, wherein the sodium blocking agent is 4, 7, 13, 16, 21-pentaoxa-1,10-diazabicyclo[8.8.5]-tricosane.
  - 53. (original): The method of claim 51, wherein the sample is a biological sample.

7

54. (original): The method of claim 53, wherein the biological sample is a blood sample.

sd-448989

55. (original): The method of claim 54, wherein the blood sample is a plasma, serum, red blood cell, or whole blood sample.

#### 56-57. (canceled)

- 58. (original): The method of claim 51, wherein the amount of AMP formed is inversely correlated to the amount of lithium ions in the sample.
- 59. (original): The method of claim 51, which is used in prognosis or diagnosis of a disease or disorder.
- 60. (currently amended): A method for assaying for lithium ions in a sample, which method comprises:
- a) contacting the sample with a first composition comprising adenosine 3',5'-bisphosphate (PAP);
- b) contacting the sample with a second composition comprising <u>a</u> the chimeric protein of claim 1, comprising, from N-terminus to C-terminus:
  - (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
  - (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution that retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
  - (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a lithium-sensitive 3'(2'),5'-bisphosphate nucleotidase; and

c) assessing the production of a detectable product to determine the presence or absence of lithium ions in the sample.

61. (original): The method of claim 60 further comprising first contacting the sample with a sodium blocking agent.

- 62. (original): The method of claim 61, wherein the sodium blocking agent is 4, 7, 13, 16, 21-pentaoxa-1,10-diazabicyclo[8.8.5]-tricosane.
  - 63. (original): The method of claim 60, wherein the sample is a biological sample.
  - 64. (original): The method of claim 63, wherein the biological sample is a blood sample.
- 65. (original): The method of claim 64, wherein the blood sample is a plasma, serum, red blood cell, or whole blood sample.
  - 66. (canceled)
- 67. (original): The method of claim 60, wherein the first composition further comprises 4-aminoantipyrine (4-AA), N-ethyl-N-(2-hydroxy-3-sulfopropyl)-3-m-toluidine (EHSPT), purine nucleoside phosphorylase, xanthine oxidase, and peroxidase, and the second composition further comprises adenosine deaminase, 5'-nucleotidase, and MgCl<sub>2</sub>.

9

sd-448989

68. (currently amended): A kit for assaying for lithium ion in a sample, which kit comprises:

- a) a first composition comprising <u>a</u> the chimeric protein <del>of claim 1, comprising, from N-terminus to C-terminus:</del>
  - (i) a first peptidyl fragment comprising a bacterial leader sequence as set forth in SEQ ID NO:1;
  - (ii) a second peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:2 or a derivative thereof having a conservative amino acid substitution that retains at least 50% of the 3'(2'),5'-bisphosphonate activity of SEQ ID NO:2; and
  - (iii) a third peptidyl fragment comprising an amino acid sequence as set forth in SEQ ID NO:3;

wherein the chimeric protein comprises comprising a lithium-sensitive 3'(2'),5'-bisphosphate nucleotidase; and

- b) a means for assessing the adenosine 3',5'-bisphosphate consumed or the AMP or Pi formed by the 3'(2'),5'-bisphosphate nucleotidase to determine the presence or amount of said lithium ions in the sample.
- 69. (previously presented): The kit of claim 68 further comprising a sodium blocking agent.
- 70. (original): The kit of claim 68, wherein the first composition further comprises adenosine deaminase, 5'-nucleotidase and MgCl<sub>2</sub>.
- 71. (previously presented): The kit of claim 68, further comprising a second composition comprising 4-aminoantipyrine (4-AA), N-ethyl-N-(2-hydroxy-3-sulfopropyl)-3-m-toluidine (EHSPT), purine nucleoside phosphorylase, xanthine oxidase, and peroxidase, wherein the reaction of 4-AA and EHSPT in the presence of

sd-448989 10

peroxidase is the means for assessing the product formed.

72. (original): The kit of claim 68, which further comprises a low lithium serum standard, a medium lithium sodium standard, and a high lithium serum standard.